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RAW SEQUENCE LISTING

Input Set : A:\Reisnerl.app

PATENT APPLICATION: US/09/202,181

DATE: 07/12/2000 TIME: 13:45:01

288

327

ENTERED

Output Set: N:\CRF3\07122000\I202181.raw 3 <110> APPLICANT: REISNER, Yair et al. 5 <120> TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO THE HEPATITIS B SURFACE ANTIGEN 8 <130> FILE REFERENCE: REISNER=5 10 <140> CURRENT APPLICATION NUMBER: 09/202,181 11 <141> CURRENT FILING DATE: 1998-12-10 13 <150> PRIOR APPLICATION NUMBER: 118625 14 <151> PRIOR FILING DATE: 1996-06-11 16 <150> PRIOR APPLICATION NUMBER: IL97/00184 17 <151> PRIOR FILING DATE: 1997-06-10 19 <160> NUMBER OF SEQ ID NOS: 4 21 <170> SOFTWARE: PatentIn Ver. 2.1 23 <210> SEQ ID NO: 1 24 <211> LENGTH: 327 25 <212> TYPE: DNA 26 <213> ORGANISM: human 28 <220> FEATURE: 29 <221> NAME/KEY: CDS 30 <222> LOCATION: (1)..(327) 32 <400> SEQUENCE: 1 33 tee tat gtg etg aet eag eea eee teg gtg tea gtg gee eea gga aag 34 Ser Tyr Val Leu Thr Gln Pro Pro Ser Val Ser Val Ala Pro Gly Lys 5 10 37 acg gcc agg att tcc tgt ggg gga aac aac att gga act aaa aat gtg 38 Thr Ala Arg Ile Ser Cys Gly Gly Asn Asn Ile Gly Thr Lys Asn Val 20 25 30 41 cac tgg tac cag cag aag cca ggc cag gcc cct gtg ctg gtc gtc tat 42 His Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Val Tyr 40 45 gct gat agc gac cgg ccc tca ggg atc cct gag cga ttc tct ggc tcc 46 Ala Asp Ser Asp Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser 47 50 55 60 49 aac tot ggg aac acg gcc acc ctg acc atc agc agg gtc gaa gtc ggg 240 50 Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Arg Val Glu Val Gly 75 51 65 70 80

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64 <212> TYPE: PRT

65 <213> ORGANISM: human

67 <400> SEQUENCE: 2

68 Ser Tyr Val Leu Thr Gln Pro Pro Ser Val Ser Val Ala Pro Gly Lys

53 gat gag gcc gac tat tac tgt cag gtg tgg gat agt gtt agt tat cat

54 Asp Glu Ala Asp Tyr Tyr Cys Gln Val Trp Asp Ser Val Ser Tyr His

57 gtg gta ttt ggc gga ggg acc acg ctg acc gtc cta ggt

58 Val Val Phe Gly Gly Gly Thr Thr Leu Thr Val Leu Gly

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71 Thr Ala Arg Ile Ser Cys Gly Gly Asn Asn Ile Gly Thr Lys Asn Val 72 20 25 30
74 His Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Val Tyr
75 35 40 45
77 Ala Asp Ser Asp Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser 78 50 60
80 Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Arg Val Glu Val Gly
81 65 70 75 80
83 Asp Glu Ala Asp Tyr Tyr Cys Gln Val Trp Asp Ser Val Ser Tyr His
84 85 90 95
86 Val Val Phe Gly Gly Gly Thr Thr Leu Thr Val Leu Gly 87 100 105
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92 <211> LENGTH: 363
93 <212> TYPE: DNA
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105 tcc ctg aga ctc tcc tgt gca ccg tct gga ttc gtc ttc agg agt tat 106 Ser Leu Arg Leu Ser Cys Ala Pro Ser Gly Phe Val Phe Arg Ser Tyr 107 20 25 30
109 ggc atg cac tgg gtc cgc cag act cca ggc aag ggg ctt gag tgg gtg
110 Gly Met His Trp Val Arg Gln Thr Pro Gly Lys Gly Leu Glu Trp Val
111 35
                                     40
                                                               45
113 tca ctt ata tgg cat gat gga agt aat aga ttc tat gca gac tcc gtg
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114 Ser Leu Ile Trp His Asp Gly Ser Asn Arg Phe Tyr Ala Asp Ser Val
115 50 55 60
117 aag ggc cga ttc acc atc tcc aga gac aat tcc aag aac aca ttg tat
118 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
119 65 70 80
121 ttg caa atg aac agc ctg aga gcc gaa gac acg gct atg tac ttc tgt
122 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Met Tyr Phe Cys
123 85 90 95
                                                                                       288
125 gcg aga gag agg ctg att gca gca cct gct gcc ttt gac ctc tgg ggc
                                                                                       336
126 Ala Arg Glu Arg Leu Ile Ala Ala Pro Ala Ala Phe Asp Leu Trp Gly
127 100 105 110
129 cag gga acc ctg gtc acc gtc tcc tcg
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136 <212> TYPE: PRT
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VERIFICATION SUMMARY
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